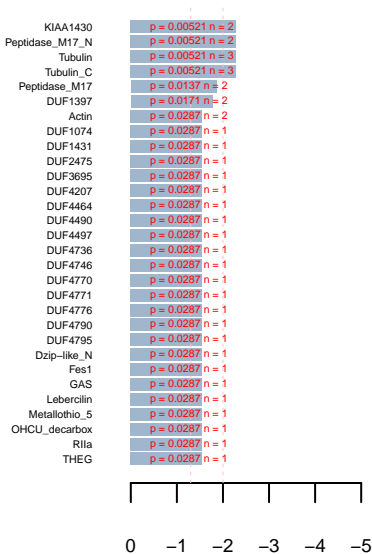


Oecy.Fem-Mal_difneg domain



log(p)

n = 144 genes | p adjust BH TRUE